

Serial Number: 09/913,419CRF Processing Date: 4/3/2003Edited by: [Signature]Verified by: [Signature] (STIC staff)

# 11

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☒ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: 3
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☐ Deleted: ☐ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file;  
☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.



OIPF

## RAW SEQUENCE LISTING

DATE: 04/03/2003

PATENT APPLICATION: US/09/913,419

TIME: 13:56:47

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\04032003\I913419.raw

3 <110> APPLICANT: Medvet Science Pty Ltd  
 4 Bagley, Christopher J  
 5 Rossjohn, Jamie  
 6 McKinsty, William J  
 7 Woodcock, Joanna M  
 8 Parker, Michael W  
 9 Lopez, Angel F  
 11 <120> TITLE OF INVENTION: CYTOKINE-BINDING DOMAIN  
 13 <130> FILE REFERENCE: 3991/0J678-US0  
 15 <140> CURRENT APPLICATION NUMBER: US 09/913,419  
 16 <141> CURRENT FILING DATE: 2001-08-08  
 18 <150> PRIOR APPLICATION NUMBER: PCT/AU00/00079  
 19 <151> PRIOR FILING DATE: 2000-02-08  
 21 <150> PRIOR APPLICATION NUMBER: PP 8576  
 22 <151> PRIOR FILING DATE: 1999-02-08  
 24 <150> PRIOR APPLICATION NUMBER: PP 8577  
 25 <151> PRIOR FILING DATE: 1999-02-09  
 27 <150> PRIOR APPLICATION NUMBER: PQ 0264  
 28 <151> PRIOR FILING DATE: 1999-05-11  
 30 <160> NUMBER OF SEQ ID NOS: 3  
 32 <170> SOFTWARE: PatentIn version 3.1  
 34 <210> SEQ ID NO: 1  
 35 <211> LENGTH: 5  
 36 <212> TYPE: PRT  
 37 <213> ORGANISM: Homo sapiens  
 39 <220> FEATURE:  
 40 <221> NAME/KEY: MISC\_FEATURE  
 41 <222> LOCATION: (3)..(3)  
 42 <223> OTHER INFORMATION: where X is any residue  
 45 <400> SEQUENCE: 1

W--&gt; 47 Trp Ser Xaa Trp Ser

48 1 5  
 51 <210> SEQ ID NO: 2  
 52 <211> LENGTH: 4  
 53 <212> TYPE: PRT  
 54 <213> ORGANISM: Homo sapiens  
 56 <220> FEATURE:  
 57 <221> NAME/KEY: MISC\_FEATURE  
 58 <222> LOCATION: (365)..(368)  
 59 <223> OTHER INFORMATION: segment of mutant cytokine binding domain  
 62 <400> SEQUENCE: 2  
 64 Ala Ala Ala Ala  
 65 1

## RAW SEQUENCE LISTING

DATE: 04/03/2003

PATENT APPLICATION: US/09/913,419

TIME: 13:56:47

Input Set : A:\PTO.AMC.txt

Output Set: N:\CRF4\04032003\I913419.raw

68 <210> SEQ ID NO: 3  
69 <211> LENGTH: 5  
70 <212> TYPE: PRT  
71 <213> ORGANISM: Homo sapiens  
73 <400> SEQUENCE: 3  
75 Trp Ser Glu Trp Ser  
76 1 5

RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/913,419

DATE: 04/03/2003  
TIME: 13:56:48

Input Set : A:\PTO.AMC.txt  
Output Set: N:\CRF4\04032003\I913419.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; Xaa Pos. 3



OIPE

## RAW SEQUENCE LISTING

DATE: 03/31/2003

PATENT APPLICATION: US/09/913,419

TIME: 11:12:46

Input Set : A:\Sequence Listing (ASCII copy).txt

Output Set: N:\CRF4\03312003\I913419.raw

**Does Not Comply  
Corrected Diskette Needed**

3 <110> APPLICANT: Medvet Science Pty Ltd  
 4 Bagley, Christopher J  
 5 Rossjohn, Jamie  
 6 McKinstry, William J  
 7 Woodcock, Joanna M  
 8 Parker, Michael W  
 9 Lopez, Angel F  
 11 <120> TITLE OF INVENTION: CYTOKINE-BINDING DOMAIN  
 13 <130> FILE REFERENCE: 3991/0J678-US0  
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 18 <150> PRIOR APPLICATION NUMBER: PCT/AU00/00079  
 19 <151> PRIOR FILING DATE: 2000-02-08  
 21 <150> PRIOR APPLICATION NUMBER: PP 8576  
 22 <151> PRIOR FILING DATE: 1999-02-08  
 24 <150> PRIOR APPLICATION NUMBER: PP 8577  
 25 <151> PRIOR FILING DATE: 1999-02-09  
 27 <150> PRIOR APPLICATION NUMBER: PQ 0264  
 28 <151> PRIOR FILING DATE: 1999-05-11  
 30 <160> NUMBER OF SEQ ID NOS: 3  
 32 <170> SOFTWARE: PatentIn version 3.1

## ERRORED SEQUENCES

68 <210> SEQ ID NO: 3  
 69 <211> LENGTH: 5  
 70 <212> TYPE: PRT  
 71 <213> ORGANISM: Homo sapiens  
 E--> 73 <400> SEQUENCE: 13  
 75 Trp Ser Glu Trp Ser  
 76 1 5

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/913,419

DATE: 03/31/2003

TIME: 11:12:47

Input Set : A:\Sequence Listing (ASCII copy).txt

Output Set: N:\CRF4\03312003\I913419.raw

L:47 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:1 after pos.:0

L:73 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:3 differs:1